

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 148592

TO: Vanessa L Ford Location: 3b25 / 3c18 Monday, March 28, 2005

Art Unit: 1645

Phone: 571-272-0857

Serial Number: 10 / 017168

From: Jan Delaval

Location: Biotech-Chem Library

Remsen 1a51

Phone: 571-272-22504

jan.delaval@uspto.gov

Search Notes	



From:

Chan, Christina

Sent:

Wednesday, March 23, 2005 8:02 AM Ford, Vanessa; STIC-Biotech/ChemLib RE: In re: 10/017,168 Sequence search

Subject:

Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89

-----Original Message-----

From:

Ford, Vanessa

Sent:

Tuesday, March 22, 2005 7:04 PM

To:

Chan, Christina

Subject:

In re: 10/017,168 Sequence search

Please search SEQ ID NOs: 1, 3,5,19, 21, 23 and 25. Please include interference searches. Please rush.

Vanessa L. Ford

Biotechnology Patent Examiner Office: REM 3B25

Mailbox: REM 3C18 Phone: 571.272.0857

Art unit:1645

STAFF USE ONLY Searcher: Searcher Phone: 2- でいく Date Searcher Picked up: 3123 Date Completed: 3/26/6 Searcher Prep/Rev. Time; Online Time:___

Type of Search

NA#:___ Interference: Oligomer:_ S/L:_ Encode/Transl:

Text: Structure#: Litigation:___ Inventor:___

Vendors and cost where applicable STN:_ DIALOG: QUESTEL/ORBIT: LEXIS/NEXIS:_

SEQUENCE SYSTEM:_ WWW/Internet:_ Other(Specify):_

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Aav85195 T. pallid
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RESULT 1 AAC84647

AAC84647 standard; DNA; 2945 BP.

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Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological sample.
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P-PSDB; AAB48316.
                                                                                                              Liu H,
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                                                                                                                                                                                                                                                                  Treponema pallidum; ssp. pallidum.
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20-APR-2001
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The invention relates to a method of detecting presence of Treponema

Claim 19; Fig 5; 73pp; English.

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CB905393 tric074xd

CB905393 tric074xd

CG744520 P037-1-E1

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Best Local Similarity 51.4%;
Matches 436; Conservative
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1 (bases 1 to 1484)
Kremitzki,C., Carter,J., McPherson,J., Warren,W., Graves,T., Mardis,E. and Wilson,R.
A physical map of the xenopus tropicalis genome
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Washington University School of Medicine
Email: submissions@watson.wustl.edu
Insert Length: 175000 Std Error: 0.00
Seq primer: Sp5 acctgccgtttcgatcct
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	(Nichols strain) " /codon start=1 /codon start=1 /trans[table=11 /trans[table=11 /protein id="CAC24646.1" /brotein id="CAC24646.1" /db xref="GI:12329841" /db	/note="unnamed protein product; Subspecies: pallidum	/mol_type="unassigned DNA" /db_xref="taxon:160" q192217		1 20AAT	THE GOVERNMENT OF THE UNITED STATES OF AMERICA (US)	}	Compositions and methods for detecting Treponema palliqum	Liu, H., Steiner, B. and Rhodes, B.	P	Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema.	Treponema pallidum		AX068032.1 GI:12329840		1 from Patent W	AX068032 2945 bp DNA linear PAT 21-JAN-2001	

1 GTCGATGCACAGCTGACGCTCTCACGTCTTGCACATATTGCGCGGGGTGCCGACATCT 60

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	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5	10.5
	10774	10596	10596	10596	10561	10516	10477	10380	10330	10285	10285	9600	9482	8705	8705	5452	2580	2580	1926	1926	1926	1925	799	795
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ALIGNMENTS	ADL67153	AAT40348	AAX15650	AAQ51731	ADL67148	ADL67150	ADL67152	AAZ22248	ADL67154	ABS66453	ABS71027	AAV21683	ADP64415	ADM10659	AAZ23778	AAX90923	AAI64275	AAA75454	ADK65580	AAF82902	AAA50254	AAX90924	AAV55831	AAV55830
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RESULT 1
AAC84648
Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological
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                                                                                                                                                                                                                                                                                                                 T. pallidum (CDC-2) acidic repeat protein (arp) encoding DNA
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P-PSDB; AAB48317.
                                                                                                              Liu H,
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20-APR-2001
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The invention relates to a method of detecting presence of Treponema

Claim 19; Fig 7; 73pp; English.

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Copyright (c) 1993 - 2005 Compugen
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CL836578 OR CBA006
CY254879 WSO2411.B
AG562801 MUS MUSCUL
AZ337339 1M0068B03
AL099791 Drosophil
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AZ334000 1M0063P17
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68.4	68.4	68.4	68.4	68.4	68.4	68.4	68.4	68.4	68.4	68.6	68.6	68.6	68.6	69	. 69	69.2	69.2	69.4	69.4	70
9.8	9.8	9.8	9.8	9.8	9.8		9.8		9.8		9.8			9.9	9.9	9.9	9.9	9.9	9.9	10.0
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CV098193	CV097413	CV093263	CV096641	CV092434	CV099895	CV092833	CN605345	CV096855	CV094178	AG435591	BQ847872	CF607575	CF609053	AG390365	BM900946	CL492192	CD493768	BZ051981	CK286024	AG346705
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REFERENCE
AUTHORS
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AUTHORS
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Query Match
Best Local Similarity
Matches 155; Conserv
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JOURNAL
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                                                                                                                                                                                                                                                                                                         alicat@sanger.ac.uk
see http://www.ebi.ac.uk/parasites/leish.html
betalls of Leishmania sequencing at the Sanger Centre are
at http://www.sanger.ac.uk/Projects/L major/
The cLHYG t3Hyg primer sequence can be obtained from acc.
                                                                                                                                                                                                                                                                                                                                                                                                                     2 (bases 1 to 67)
Taylor,R.G., Huckle,E.E.J., Ivens,A.C., Rajandream,M.A. and
Taylor,R.G.,
Barrell,B.G.
Direct Submission
Submitted (14-MAR-2000) Leishmania major Friedlin genome sequencing
project Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 365)
Ivens,A.C., Lewis,S.M., Bagherzadeh,A., Zhang,L., Chan,H.M.
Smith,D.F.
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L2644X

Leishmaia major Friedlin cosmid L2644 tJHyg end-sequence, smilar to U40394 U40394 Kaposi's sarcoma-associated herpes-like..., N=755, Prob=1.8e-54; TR:P93797 P93797 PHEROPHORIN-S PRECURSOR. [0] +3..., N=318, Prob=7.2e-37, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A physical map of the Leishmania major Friedlin genome Genome Res. 8 (2), 135-145 (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Leishmania major
Leishmania major
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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12.8%;
nilarity 58.7%;
Conservative (
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/mol type="genomic DNA"
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/chromosome="26"
/clone="cosmid L2644"
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Score 89.6; DB 9;
Pred. No. 4.5e-13;
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              693.4
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12.6 265556
12.5 233939
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10846.472 Million cell updates/sec
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AX068036 Sequence
AF342807 Treponema
AF4111124 Treponema
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AC121566 Mus muscu
AC121566 Mus muscu
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Qy 1 ATC	Query Match Best Local Similarity Matches 699; Conser	ORIGIN	CDS	AUTHORS LIU,H., AUTHORS LIU,H., TITLE Composi JOURNAL Patent: THE GOV FEATURES SOURCE	Z Z	AX068034
ATGTTTGTGCGCAGTGACATGTTCCCCAAAAACACTGCTGTTGAAATTAGCAACTTAGAA 	100.0%; Score 699; DB (100.0%; Pred. No. 2.5e-1 vative 0; Mismatches	/trans1 table=11 /protein_id="CAC24647.1" /protein_id="CAC24647.1" /db_xref="G1:1229843" /db_xref="G1:1229843" /trans1ation="MFVRSDMFPKNTAVEISNLEKNAKAQAVVIGHAGIPGLLVSLAP /trans1ation="MFVRSDMFPKNTAVEISNLEKNAKAQAVVIGHAGIPGLLVSLAP AAAAQLGIGVYQAVRVRTLGTVRGGSQTSQDGLSLASLPSRVPARPAQRDPLSSPP AGHTVPEYRDTVIFDDPRLVSPLSREVEDVPKVVEPAJSREGGEREVEDVPKVVEPAS EREGGEREVEDVPKVVEPASEREGGEREVEDVPKVVE:"ASEREGGEREVASQHTKQPS HSVSNSAPNQFRKP"	/organism="Treponema pallidum" /mol_type="unassigned DNA" /db_xref="taxon:160" 1699 1699 /note="unnamed protein product; (CDC-2 strain)"	T WC	AXO68034 699 bp DNA Sequence 3 from Patent WO0077486. AXO68034 AXO68034.1 GI:12329842 Treponema pallidum Treponema pallidum Treponema pallidum Treponema pallidum	
CTGTTGAAATTAGCAACTTAGAA 60	; Length 699; 145; 0; Indels 0; Gaps 0;	SNLEKNAKAQAVVIGHAGIPGLLVSLAP SQDGLSLASLPSRVPARPAQRDPLSSPP PKVVEPASEREGGEREVEDVPKVVEPAS DVPKVVE?ASEREGGEREVASQHTKQPS	; Subspecies: pertenue	Treponema pallidum AMERICA (US)	linear PAT 19-JAN-2001	

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18.0 211580
17.4 19741
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16.2 216602
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AC117639	AC123837	AL591208	AC121865	AC125192	AC122889	AC102602	AC121566	AC084389	AC098712	AC126271	AC125235	AC122232	AC122052	AC103637	AC122851	AC123857	AC120236_0	AC109193	AC124108	AC091464	AC117585	BX005304	AC118476	AC127583	AC098271	
Mus	AC123837 Mus muscu	AL591208 Mouse DNA	AC121865 Mus muscu	Mus	AC122889 Mus muscu	AC102602 Mus muscu	AC121566 Mus muscu	AC084389 Mus muscu			AC125235 Mus muscu	AC122232 Mus muscu	Mus	Mug	AC122851 Mus muscu	AC123857 Mus muscu	AC120236 Rattus no	AC109193 Mus muscu	Mus n	AC091464 Mus muscu	AC117585 Mus muscu		AC118476 Mus muscu	AC127583 Mus muscu	AC098271 Rattus no	

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AUTHORS
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                                                          1 ATGTTTGTGCGCAGTGACATGTTCCCCCAAAAACACTGCTGTTGAAATTAGCAACTTAGAA
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Compositions and methods for detecting Treponema pallidum
Patent: WO 0077486-A 5 21-DEC-2000;
THE GOVERNMENT OF THE UNITED STATES OF AMERICA (US)
Location/Qualifiers
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Sequence 5 from Patent WO0077486.
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Treponema pallidum
Bacteria; Spirochaetes; Spirochaetales; Spirochaetaceae; Treponema.
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Aac28607 T. pallid
Aac20607 polynucle
Aac84648 T. pallid
Aac20608 Nouse can
Aav5831 Nucleotid
Aaa50254 Epstein B
Aaf82902 EBV tethe
Ad645580 Human her
Aaa77454 Nucleotid
Aai6475 Epstein B
Aax90923 Anti-sen pS
Adm10659 Expressio
Adm10659 Expressio
Adm264415 Vector pC
Aav21683 Vector pl
Abs71107 pCEP-Xa-F
Abs66453 Plasmid p
Ad167154 Plasmid p
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12.7	12.7	12.7	12.9	13.3		8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8	13.8
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Abk98592 Vector pE	Acd13882 L. lactis	Abk98631 Vector pE	Adq97246 Mouse can	Ach89664 Human gen	Aav55830 FLGA inse	Aax90924 Epstein B	Adn12161 Epstein-B	Adl71910 Expressio	Aaa59553 DNA clone	Ado07394 Modified	Ado07395 Modified	~	Adl67176 Plasmid p	Adl67147 Plasmid p	Adl67149 Plasmid p	Adl67151 Plasmid p	Adl67153 Plasmid p	Adl67175 Plasmid p	Aat40348 Plasmid p	Nucleot	Plasmid p	Adl67148 Plasmid p	Adl67150 Plasmid p	Adl67152 Plasmid p

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Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological sample.
                                                                                                                                                                                                                                                        Treponema pallidum; acidic repeat protein; arp; immunogenic; syphilis; yaws; bejel; ds.
                                                                                                                                                                                                                                                                                  T. pallidum (Bosnia) acidic repeat protein (arp) encoding DNA.
                                                                                                                                                                                                                                       Treponema pallidum; ssp. endemicum
                                                                                                                     (USSH ) US DEPT HEALTH & HUMAN SERVICES.
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Aaz22248 Nucleotid

The invention relates to a method of detecting presence of Treponema

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Copyright (c) 1993 - 2005 Compugen Ltd.
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12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.6	12.7	12.7	12.7	12.7	12.7
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A:3349830 Mus muscu	C2877292 tric075xf	CB905602 tric075xf	CC578225 CH240_457	AG382169 Mus muscu	CG748176 P042-1-A0	AG448606 Mus muscu	CR116264 Forward B	A2186646 SP_1007_B	CR074624 Forward s	BX121682 Danio rer	AG435617 Mus muscu	CI/080508 CH216-158	CC300639 CH261-68J	CI.086207 ISB1-7A17	CC298943 CH261-22C	AG396109 Mus muscu	CC262597 CH261-167	AL453768 T. brucei	CC294165 CH261-130	BI958090 HVSMEn001

	SOUTCE	FEATURES											COMMENT				JOURNAL	TITLE	AUTHORS	REFERENCE	JOURNAL	TITLE	AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	AG561854/c	RESULT 1
/organism="Mus musculus molossinus" /organism="Mus musculus molossinus" /mol type="genomic DNA" /sub_species="molossinus" /db xref="taxon:57486" /clone="MSMg01-481H05.T7" /sex="male" /tissue type="mixture of kidney and spleen" /clone_Tib="MSMg01 Mouse Male BAC Library"	1 776	Location/Qualifiers		•	¥	Sequencing: T7	PRIMERS	e-mail: abe@rtc.riken.jp	phone: 81-298-36-9189, fax: 81-298-36-9199	Koyadai, Tsukuba, 305-0074 Japan	Tsukuba Institude, Bio Resource Center,	library availability, please contact Kuniya Abe (abe@rtc.riken.jp).	Clones are derived from the mouse BAC library MSMg01. For BAC	Tel:81-45-503-9111, Fax:81-45-503-9170)	(E-mail:hattori@gsc.riken.jp, URL:http://hgp.gsc.riken.go.jp/,	1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan	Submitted (17-NOV-2003) Masahira Hattori, The Institute of Physical	ission	Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y.	2 (bases 1 to 776)	Unpublished		Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y.	1	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.		Mus musculus molossinus	GSS.	AG561854.1 GI:48322552	AG561854	COMMONDALINA DNA, CIONE: MONGOLITA CIONOCIA, SENONIA GUIVEZ	1		

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10846.472 Million cell updates/sec
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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289751	199854	253504	165592	125681	199459	266556	215769	178757	211580	699	939	2946	2945	1647	1047	1287	14268	1182	Match Length	
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AC091458	AL591208	AC102602	AC122020	AC122889	AC125192	AC136868	AC100209	AC122509	AC111089	AX068034	AX068036	AF015824	AX068032	AF411124	AF342806	AF342807	AE001220	AF411126	ij	
	AL591208 Mouse DNA	AC102602 Mus muscu	AC122020 Mus muscu	AC122889 Mus muscu	AC125192 Mus muscu	AC136868 Rattus no	AC100209 Mus muscu	AC122509 Mus muscu	AC111089 Mus muscu	. AX068034 Sequence	AX068036 Sequence	AF015824 Treponema	AX068032 Sequence	AF411124 Treponema	AF342806 Treponema	AF342807 Treponema	AE001220 Treponema	AF411126 Treponema	Description	

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10.1	10.1	10.1	10.1	10.1	10.1	10.1	10.1	10.1	10.2	10.2	10.2	10.3	10.3	10.3	10.3	10.3	10.3	10.3	10.3	10.3	10.3	10.4	10.4	10.4
182777	172853	211161	215745	203086	197683	196236	191753	154462	318930	110000	163945	208880	182272	260625	229190	219771	217353	216602	171593	120487	178258	139630	197411	152435
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AC115928 AC123857	AC120147	AC098271	AC117585	AC099948	AC118476	AC108780	AC125235	AC145149	AC073495	AE016753_2	AC116727	AC123837	AC122232	AC096018	AC127348	BX005376	AC149611	AC136729	AC122052	AC103637	AC109296	AC133945	AC135961	AC121566
AC115928 Mus muscu AC123857 Mus muscu	AC120147 Mus muscu	AC098271 Rattus no	AC117585 Mus muscu	AC099948 Mus muscu	AC118476 Mus muscu	AC108780 Mus muscu	AC125235 Mus muscu	AC145149 Homo sapi	AC073495 Mus muscu	Continuation (3 of	AC116727 Mus muscu	AC123837 Mus muscu	AC122232 Mus muscu	AC096018 Rattus no	AC127348 Mus muscu	BX005376 Danio rer	AC149611 Mus muscu	AC136729 Mus muscu	FC122052 Mus muscu	A.C103637 Mus muscu	AC109296 Mus muscu	AC133945 Mus muscu	AC135961 Mus muscu	AC121566 Mus muscu

	CDS	gene			source	FEATURES		JOURNAL	TITLE	AUTHORS	REFERENCE	JOURNAL		TITLE	AUTHORS	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	FOCUS	AF#IIIZ6	RESULT 1	1
/gene="atp" /codon start=1 /codon start=1 /transT_table=11 /transT_table=11 /product="acidic repeat protein" /product="acidic repeat protein" /protein id="akl07373.1" /db_xref="G1:15778315" /db_xref="G1:15778315" /db_xref="G1:15778315" /db_xref="G1:15778315" /db_xref="G1:15778315" /db_xref="G1:15778315" /db_xref="G1:15778315" /db_xref="G1:15778315" /protein id="akl07373.1" /db_xref="G1:15778315" /db_xref="G1:15778315" /protein id="akl07373.1" /db_xref="G1:15778315" /db_xref="G1:15778315" /protein id="akl07355" /db_xref="G1:15778315" /db_xref="G1:15778315" /protein id="akl07355" /db_xref="G1:15778315" /db_xref="G1:15778315" /protein" /prot	1. 1182	11182	/sub_species="pertenue" /db_xref="taxon:168"	,	/organisms"Trenonema pallidum subsp. mertenue"	Location/Qualifiers	Centers for Disease Control and Prevention, 1600 Clifton Rd, Atlanta, GA 30333, USA		Direct Submission	Liu, H., Steiner, B.M. and Rodes, B.	2 (bases 1 to 1182)	Unpublished	of Treponema pallidum	Molecular characterization of the acidic repeat protein gene (arp)	Liu, H., Steiner, B.M. and Rodes, B.	bacteria; spirochaetes; spirochaetates; spirochaetateae; ireponema. 1 (bases 1 to 1182)	pallidum subsp. pertenue			AF411126.1 GI:15778314	AP411126	protein (arm) gene, complete cds.	DNA			

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	FEATURES BOUTCE	COMMENT	AUTHORS TITLE JOURNAL REFERENCE AUTHORS TITLE TITLE JOURNAL	ACCESSION VERSION KEYWORDS SOURCE ORGANISM	RESULT 1 AG561854/c LOCUS DEFINITION
/organism="Mus musculus molossinus" /mol_type="genomic DNA" /sub_species="molossinus" /db_xref="taxon:57486" /clone="MSMg01-481H05.T7" /sex="male" /tissue_type="mixture_of_kidney_and_spleen" /clone_Tib="MSMg01 Mouse_Male_BAC_Library"	PRIMERS Sequencing: T7 LIBRARY Vector: pBACe3.6 R.Site 1: EcoRI R.Site 2: EcoRI Location/Qualifiers 1. 776	mical Research (RIKEN), Genomic Sciences Center (GSC Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045. hattoriegsc.riken.jp, URL:http://hpp.gsc.riken.go.:45-503-9111, Fax:81-45-503-9170) are derived from the mouse BAC library MSMg01. For It availability, please contact Kuniya Abe (abe@rtc.ri. Institude, Bio Resource Center, titute of Physical and Chemical Research (RIKEN) 3-17 Sukuba, 305-0074 Japan 81-298-36-9189, fax: 81-298-36-9199 abe@rtc.riken.jp	Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y. BAC end Sequences of Library MSMg01 Unpublished 2 (bases 1 to 776) Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y. Direct Submission Submitted (17-NOV-2003) Masahira Hattori, The Institute of Physical	Bequence. AG561854 AG561854.1 GI:48322552 GSS. Mus musculus molossinus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.	AG561854 776 bp DNA linear GSS 05-JUN-2004 Mus musculus molossinus DNA, clone:MSMg01-481H05.T7, genomic survey

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8.3	8.3	8.5	8.8				9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0	9.0
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RESULT 1
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DT 05-N
XX Treg
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AAX20500-21243 represent polynucleotide sequences from the genome of Treponema pallidum. The sequences can be used for detection, diagnosis, characterisation, prevention and therapy for T. pallidum infections, particularly syphilis. They can also be used for detecting diseases related to Borrelia infections in animals, and for the production of

Sequence 9410 BP; 1934 A; 2470 C; 2830 G; biosynthetic products such as enzymes

2153 T;

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23 Other;

New isolated Treponema pallidum nucleic acids - used to develop products for the detection, diagnosis, characterisation, prevention and therapy of T. pallidum infections, particularly syphilis.

WPI; 1999-081273/07.

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Claim 1; Page 693-698; 1150pp; English.

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Copyright (c) 1993 - 2005 Compugen Ltd.
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10	169.4	13.2	211580	N	AC111089	AC111089 Mus muscu
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12	154.4	12.0	197411	10	AC135961	AC135961 Mus muscu
c 13	154	12.0	260625	N	AC096018	AC096018 Rattus no
c 14	150.2	11.7	318930	N	AC073495	AC073495 Mus muscu
c 15	149.6	11.6	216602	10	AC136729	AC136729 Mus muscu
c 16	148.6	11.5	163945	N	AC116727	AC116727 Mus muscu
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/organisms"ireponema Pallium subsp. endemicum /mol type="genomic DNA" /strain="Bosnia" /sub_species="endemicum" /db_xref="taxon:53436"			On Sep 13, 2001 this sequence version replaced gi:12667500.	Stop D13, Atlanta, GA 30333, USA	and Prevention,	-2001) National Center for	Direct Submission	Liu, H., Steiner, B. and Rodes, B.	4 (bases 1 to 1287)	on i	Submitted (25-JAN-2001) National Center for Infectious Diseases,	Direct Submission	Tin H and Atoiner B	3 (bases 1 to 1287)	Unpublished	Treponema pallidum	N	Liu.H. Steiner, B.M. and Rodes, B.	2 (bases 1 to 1287)	Unpublished	(Bosnia strain)	Acidic repeat protein (arp) gene sequence of Treponema pallidum	Liu, H. and Steiner, B.		Spirochaetes; Sp		Treponema pallidum subsp. endemicum		AF342807.2 GI:15617206		protein (arp) gene, complete cds.	Treponema pallidum subsp. endemicum strain Bosnia acidic repeat	AF342807 1287 bp DNA linear BCT 13-SEP-2001		

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ALIGNMENTS

REFERENCE AUTHORS TITLE JOURNAL REFERENCE SOURCE ORGANISM ACCESSION VERSION KEYWORDS RESULT 1 AG561854/c LOCUS COMMENT FEATURES DEFINITION JOURNAL source Vector R.Site R.Site Direct Submission Submitted (17-NOV-2003) Masshira Hattori, The Institute of Physical Submitted (17-NOV-2003) Masshira Hattori, The Institute of Physical and Chemical Research (RIKEN), Genomic Sciences Center (GSC); 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan 1-7-22 Suehiro-chou, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan 16-811-14-503-9111, Fax:881-45-503-9170) Tel:81-45-503-9111, Fax:881-45-503-9170) Clones are derived from the mouse BAC library NSMg01. For BAC (1) library availability, please contact Kuniya Abe Tsukuba Institude, Bio Resource Center, The Institute of Physical and Chemical Research Koyadai, Tsukuba, 305-0074 Japan phone: 81-298-36-9189, fax: 81-298-36-9199 Mus musculus molossinus Mus musculus molossinus Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus Mus musculus molossinus Sequencing : T7 LIBRARY Hattori,M., Toyoda,A., Noguchi,H., Kojima,T. and Sakaki,Y. BAC end Sequences of Library MSMg01 sequence. AG561854 PRIMERS e-mail: abe@rtc.riken.jp 2 (bases 1 to 776) Hattori, M., Toyoda, A., Noguchi, H., Kojima, T. and Sakaki, Y. Unpublished AG561854.1 AG561854 N 14 /organism="Mus musculus molossinus" /mol type="genomic DNA" /sub_species="molossinus" /db xref="taxon.57486" /clone="MSMg01-481H05.T7" /sex="male" Location/Qualifiers tissue_type="mixture of kidney and spleen" /clone_lib="MSMg01 Mouse Male BAC Library" : pBACe3.6 : EcoRI GI:48322552 ECORI DNA, 776 bp DNA linear (clone:MSMg01-481H05.T7, library NSMg01. For BAC uniya Abe (abe@rtc.riken.jp) (RIKEN) 3-1-1 GSS 05-JUN-2004 , genomic survey

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A Treponema pallidum; acidic repeat protein; arp; yaws; bejel; ds. AAC84649 standard; DNA; 939 BP. T. pallidum (Bosnia) acidic repeat protein (arp) encoding DNA. 11-SEP-2003 20-APR-2001 AAC84649; Claim 19; Fig 9; 73pp; English. Detecting Treponema pallidum in blood, saliva, etc., by detecting formation of a complex between immunogenic peptides of acidic repeat protein of the bacterium and an antibody present in the biological sample. Liu H, 14-JUN-1999; 14-JUN-2000; 2000WO-US016425 21-DEC-2000. WO200077486-A2. Treponema pallidum; ssp. endemicum WPI; 2001-080711/09. P-PSDB; AAB48318. (USSH) US DEPT HEALTH & HUMAN SERVICES Steiner B, (revised) (first entry) 99US-0138981P. Location/Qualifiers
1. .939 Rhodes B; immunogenic; syphilis;

The invention relates to a method of detecting presence of Treponema

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Result
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,	REMARK COMMENT FEATURES SOURCE	TITLE JOURNAL	REFERENCE AUTHORS	REFERENCE AUTHORS TITLE JOURNAL	REFERENCE AUTHORS TITLE JOURNAL	REFERENCE AUTHORS TITLE JOURNAL	KEYWORDS SOURCE ORGANISM	LOCUS DEFINITION ACCESSION VERSION
/organism="Treponema pallidum subsp. pe::tenue" /mol_type="genomic DNA" /strain="DDC: /sub_species="pertenue" /db_xref="taxon:168"	Sequence update by submitter On Sep 13, 2001 this sequence version replaced gi:12667498. Location/Qualifiers 11047	Direct Submission Submitted (13-SEP-2001) National Center for Infectious Diseases, Centers for Disease Control and Prevention, 1600 Clifton Rd. Mail Stop D13, Atlanta, GA 30333, USA		3 (bases 1 to 1047) Liu,H. and Steiner,B. Direct Submission, Submitted (25-JAN-2001) National Center for Infectious Diseases,	2 (bases 1 to 1047) Liu,H., Steiner,B.M. and Rodes,B. Molecular characterization of the acidic repeat protein (arp) of Treponema pallidum Unpublished	1 (bases 1 to 1047) Liu,H. and Steiner,B. Acidic repeat protein (arp) gene sequence of Treponema pallidum subspecies pertenue (CDC 2 strain) Unpublished	Treponema pallidum subsp. pertenue (yaws treponeme) Treponema pallidum subsp. pertenue Bacteria; Spirochaetes; Spirochaetales; Spirochaetareae. Treponema	AF342806 1047 bp DNA linear BCT 13-SEP-2001 Treponema pallidum subsp. pertenue strain CDC2 acidic repeat protein (arp) gene, complete cds. AF342806 AF342806.2 GI:15617204

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AG562801 Mus muscu
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AL0393791 Drosophi3
AL0393791 Drosophi3
AC383375 1M0141E03
CC323037 AGENCOURT
BG273097 nai99706.
CC251395 CH261-157
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<u>.</u>	Ω	편 25	
FEATURES source	TITLE JOURNAL COMMENT	REFERENCE AUTHORS TITLE JOURNAL MEDLINE PUBMED REFERENCE AUTHORS	RESULT 1 L2644X/c LOCUS DEFINITION ACCESSION VERSION KEYNORDS SOURCE ORGANISM
Location/Qualifiers 1. 365 /organism="Leishmania major" /mol_type="genomic DNA" /strain="Friedlin" /strain="Friedlin" /db_xref="taxon:5664" /chromosome="26"	Direct Submission Submitted (14-MAR-2000) Leishmania major Friedlin genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and alicat@sanger.ac.uk slicat@sanger.ac.uk see http://www.ebi.ac.uk/parasites/leish.html Details of Leishmania sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/L major/ The cLHYG t3Hyg primer sequence can be obtained from acc. no. US9231.	1 (bases 1 to 365) Ivens,A.C., Lewis,S.M., Bagherzadeh,A., Zhang,L., Chan,H.M. and Smith,D.F. Smith,D.F. A physical map of the Leishmania major Friedlin genome Genome Res. 8 (2), 135-145 (1998) 98146435 9477341 2 (bases 1 to 67) 2 (bases 1 to 67) Taylor,R.G., Huckle,E.E.J., Ivens,A.C., Rajandream,M.A. and Barrell,B.G.	L2644X L2644X L2644X L2644X L2644X L2644 L

Query Match 8.6%; Best Local Similarity 58.7%; Matches 155; Conservative

Score 89.6; DB 9; Pred. No. 1.5e-13; 0; Mismatches 109;

Length 365; Indels

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AAX20500-21243 represent polynucleotide sequences from the genome of Treponema pallidum. The sequences can be used for detection, diagnosis, characterisation, prevention at therapy for T. pallidum infections, particularly syphils. They can also be used for detecting diseases related to Borrelia infections in animals, and for the production of biosynthetic products such as enzymes

Sequence 9410 BP; 1934 A; 2470 C; 2830 G; 2153 T; 0 U; 23 Other;

New isolated Treponema pallidum nucleic acids - used to develop products for the detection, diagnosis, characterisation, prevention and therapy of T. pallidum infections, particularly syphilis.

WPI; 1999-081273/07.

Claim 1; Page 693-698; 1150pp; English.

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ALIGNMENTS

RESULT 1

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Treponema pallidum infection; syphilis; Borrelia infection; animal; enzyme production; ds.
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                                                                                                                                                                                                                                                           AAX20603 standard; DNA; 9410 BP.
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                                                                                                                                                                                                                     Polynucleotide sequence from the genome of Treponema pallidum
                                                                                                                      (HUMA-) HUMAN GENOME SCI INC
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